



## RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/807,660

Source: PCT09

Date Processed by STIC: 5-1-01

**THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.**

**PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:**

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- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

**FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.**

**FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.**

**PATENTIN 2.1 e-mail help: [patin21help@uspto.gov](mailto:patin21help@uspto.gov) or phone 703-306-4119 (R. Wax)**

**PATENTIN 3.0 e-mail help: [patin3help@uspto.gov](mailto:patin3help@uspto.gov) or phone 703-306-4119 (R. Wax)**

**TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:**

### **Checker Version 3.0**

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

**Checker Version 3.0 can be down loaded from the USPTO website at the following address:**  
**<http://www.uspto.gov/web/offices/pac/checker>**

PCT

## RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/807,660

DATE: 05/01/2001

TIME: 11:37:50

Input Set : A:\23,495 PCT.txt

Output Set: N:\CRF3\05012001\I807660.raw

Does Not Comply  
Corrected Diskette Needed  
see pp. 1, 2

3 <110> APPLICANT: Maliszewski, Charles R.  
4 Gayle III, Richard B.  
5 Marcus, Aaron J.  
6 Immunex Corporation  
7 Cornell Research Foundation, Inc.  
9 <120> TITLE OF INVENTION: Methods of Inhibiting Platelet Activation and  
10 Recruitment  
12 <130> FILE REFERENCE: 23,495 PCT  
C--> 14 <140> CURRENT APPLICATION NUMBER: US/09/807,660  
C--> 15 <141> CURRENT FILING DATE: 2001-04-16  
17 <150> PRIOR APPLICATION NUMBER: US 60/104,585  
18 <151> PRIOR FILING DATE: 1998-10-16  
20 <150> PRIOR APPLICATION NUMBER: US 60/107,466  
21 <151> PRIOR FILING DATE: 1998-11-06  
23 <150> PRIOR APPLICATION NUMBER: US 60/149,010  
24 <151> PRIOR FILING DATE: 1999-08-13  
26 <160> NUMBER OF SEQ ID NOS: 31  
28 <170> SOFTWARE: PatentIn Ver. 2.0

## ERRORED SEQUENCES

1925 <210> SEQ ID NO: 31  
1926 <211> LENGTH: 58  
1927 <212> TYPE: PRT  
1928 <213> ORGANISM: Homo sapiens  
1930 <400> SEQUENCE: 31  
1931 Met Ala Thr Ser Trp Gly Thr Val Phe Phe Met Leu Val Val Ser Cys  
1932 1 5 10 15  
1934 Val Cys Ser Ala Val Ser His Arg Asn Gln Gln Thr Trp Phe Glu Gly  
1935 20 25 30  
1937 Ile Phe Leu Ser Ser Met Cys Pro Ile Asn Val Ser Ala Ser Thr Leu  
1938 35 40 45  
1940 Tyr Gly Ile Met Phe Asp Ala Gly Ser Thr  
1941 50 55  
E--> 1945 1  
E--> 1948 1

→ Delete extraneous numerals at the end of the file. They are causing an invalid amino acid count for sequence #31.

Seg #6 (Partially shown)

<210> 6

<211> 454

<212> PRT

<213> Artificial Sequence

→ missing mandatory <220>, <223>  
feature to explain the source

<400> 6

Ala Pro Thr Ser Ser Ser Thr Lys Lys Thr Gln Leu Thr Ser Ser Thr  
1 5 10 15

of the artificial

Gln Asn Lys Ala Leu Pro Glu Asn Val Lys Tyr Gly Ile Val Leu Asp  
20 25 30

sequence.



Ala Gly Ser Ser His Thr Ser Leu Tyr Ile Tyr Lys Trp Pro Ala Glu  
35 40 45

Note: Error

Lys Glu Asn Asp Thr Gly Val Val His Gln Val Glu Glu Cys Arg Val  
50 55 60

repeated in

Sequence #5

8, 26

## VERIFICATION SUMMARY

DATE: 05/01/2001

PATENT APPLICATION: US/09/807,660

TIME: 11:37:52

Input Set : A:\23,495 PCT.txt

Output Set: N:\CRF3\05012001\I807660.raw

L:14 M:270 C: Current Application Number differs, Replaced Application Number  
L:15 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:398 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4  
L:620 M:258 W: Mandatory Feature missing, <220> FEATURE:  
L:620 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:  
L:850 M:258 W: Mandatory Feature missing, <220> FEATURE:  
L:850 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:  
L:1426 M:258 W: Mandatory Feature missing, <220> FEATURE:  
L:1426 M:258 W: Mandatory Feature missing, <223> OTHER INFORMATION:  
L:1945 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:31  
M:332 Repeated in SeqNo=31